Amendments to the Claims

Claims 1-104. (Cancelled).

105. (New) A butyrylcholinesterase variant comprising an amino acid sequence selected from the group consisting of SEQ ID NO: 4, 6, 8, 10, 12, 14, 24, 26, 28, 30, 32, 34, 36, 38, 40, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 78, 80, 82, 84, 86, 88, 90, 92, 94, 96, 98, 100, 102, 104, 106, 108, 110, 112, 114, 116, 118, 120, 122, 124, 126, 128, 130, 132, 134, 136, 138, 140, 142, 144, 146, 148, 150, 152, 154, 156, 158, 160, 162, 164, 166, 168, 170, 172, 174, 176, 178, 180, 182, 184, 186, 188, 190, 192, 194, and 196, or a functional fragment thereof, wherein the variant or fragment comprises alanine at amino acid position 227.

- 106. (New) The butyrylcholinesterase variant of Claim 105, wherein the amino acid sequence is selected from the group consisting of SEQ ID NO: 24, 26, 30, 32, 34, 36, 38, 104, 106, 108, 110, 112, 116, 118, 120, 122, 124, 126, 128, 132, 134, 136, 140, and 142, or a functional fragment thereof.
- 107. (New) The butyrylcholinesterase variant of Claim 106, wherein the amino acid sequence is selected from the group consisting of SEQ ID NO: 36, 108, 110, 112, 122, 124, 134, 178, 180, 182, 186, 188, 190, 192 and 196, or a functional fragment thereof.
- 108. (New) The butyrylcholinesterase variant of Claim 107, wherein the amino acid sequence is selected from the group consisting of SEQ ID NO: 178, 180, and 188, or a functional fragment thereof.
- 109. (New) The butyrylcholinesterase variant of Claim 108, wherein the amino acid sequence is SEQ ID NO: 180 or a functional fragment thereof.
- 110. (New) The butyrylcholinesterase variant of Claim 105, having at least a two-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 111. (New) The butyrylcholinesterase variant of Claim 110, having at least a fifty-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.

- 112. (New) The butyrylcholinesterase variant of Claim 111, having at least a one hundred-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 113. (New) The butyrylcholinesterase variant of Claim 112, having at least a five hundred-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 114. (New) The butyrylcholinesterase variant of Claim 113, having at least a fifteen hundred-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 115. (New) The butyrylcholinesterase variant of Claim 114, having at least a two thousand-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 116. (New) The butyrylcholinesterase variant of Claim 115, having at least a two thousand five hundred-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 117. (New) The butyrylcholinesterase variant of Claim 116, having at least a three thousand-fold increase in camptothecin conversion activity compared to butyrylcholinesterase, or functional fragment thereof.
- 118. (New) The butyrylcholinesterase variant of Claim 105, or functional fragment thereof, further comprising an antibody or antibody fragment which specifically binds the epidermal growth factor receptor (EGFR).
- 119. (New) The butyrylcholinesterase variant of Claim 118, wherein the antibody or antibody fragment comprises an amino acid sequence as shown in SEQ ID NO: 18 or 20.
- 120. (New) The butyrylcholinesterase variant of Claim 105, further comprising an antibody or antibody fragment which specifically binds the CD20 cell surface antigen.
- 121. (New) The butyrylcholinesterase variant of Claim 120, wherein the antibody or antibody fragment comprises an amino acid sequence as shown in SEQ ID NO: 198.
- 122. (New) A nucleic acid encoding a butyrylcholinesterase variant comprising the nucleic acid sequence selected from the group consisting of SEQ ID NO: 3, 5, 7, 9, 11, 13, 23, 25, 27, 29, 31, 33, 35, 37, 39, 41, 43, 45, 47, 49, 51, 53, 55, 57, 59, 61, 63, 65, 67, 69, 71,

- 73, 75, 77, 79, 81, 83, 85, 87, 89, 91, 93, 95, 97, 99, 101, 103, 105, 107, 109, 111, 113, 115, 117, 119, 121, 123, 125, 127, 129, 131, 133, 135, 137, 139, 141, 143, 145, 147, 149, 151, 153, 155, 157, 159, 161, 163, 165, 167, 169, 171, 173, 175, 177, 179, 181, 183, 185, 187, 189, 191, 193, and 195, or a fragment thereof.
- 123. (New) The nucleic acid of Claim 122, wherein the nucleic acid sequence is selected from the group consisting of SEQ ID NO: 177, 179, 181, 183, 185, 187, 189, 191, 193, and 195, or a fragment thereof.
- 124. (New) A method of converting a camptothecin derivative to a topoisomerase inhibitor comprising contacting said camptothecin derivative with a butyrylcholinesterase variant comprising an amino acid sequence selected from SEQ ID NOS: 2, 4, 6, 8, 10, 12, 14, 24, 26, 28, 30, 32, 34, 36, 38, 40, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 78, 80, 82, 84, 86, 88, 90, 92, 94, 96, 98, 100, 102, 104, 106, 108, 110, 112, 114, 116, 118, 120, 122, 124, 126, 128, 130, 132, 134, 136, 138, 140; 142, 144, 146, 148, 150, 152, 154, 156, 158, 160, 162, 164, 166, 168, 170, 172, 174, 176, 178, 180, 182, 184, 186, 188, 190, 192, 194 and 196, or functional fragment thereof, under conditions that allow conversion of a camptothecin derivative to a topoisomerase inhibitor.
- 125. (New) The method of Claim 124, wherein said butyrylcholinesterase variant is selected from the group consisting of SEQ ID NOS: 24, 26, 30, 32, 34, 36, 38, 104, 106, 108, 110, 112, 116, 118, 120, 122, 124, 126, 128, 132, 134, 136, 140 and 142, or functional fragment thereof.
- 126. (New) The method of Claim 125, wherein said butyrylcholinesterase variant is selected from the group consisting of SEQ ID NO: 178, 180, 182, 184, 188 and 192, or functional fragment thereof.
- 127. (New) The method of Claim 126, wherein said butyrylcholinesterase variant is SEQ ID NO: 180, or functional fragment thereof.
- 128. (New) The method of Claim 124, wherein said butyrylcholinesterase variant exhibits a two-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 129. (New) The method of Claim 128, wherein said butyrylcholinesterase variant exhibits a fifty-fold or greater increase in conversion capability compared to butyrylcholinesterase.

- 130. (New) The method of Claim 129, wherein said butyrylcholinesterase variant exhibits a one hundred-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 131. (New) The method of Claim 130, wherein said butyrylcholinesterase variant exhibits a five hundred-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 132. (New) The method of Claim 131, wherein said butyrylcholinesterase variant exhibits a fifteen hundred-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 133. (New) The method of Claim 132, wherein said butyrylcholinesterase variant exhibits a two thousand-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 134. (New) The method of Claim 133, wherein said butyrylcholinesterase variant exhibits a two thousand five hundred-fold or greater increase in conversion capability compared to butyrylcholinesterase.
- 135. (New) The method of Claim 134, wherein said butyrylcholinesterase variant exhibits a three thousand-fold or greater increase in conversion capability compared to butyrylcholinesterase.
 - 136. (New) The method of Claim 124, wherein said topoisomerase inhibitor is SN-38.
- 137. (New) The method of Claim 136, wherein said camptothecin derivative is CPT-11.
- 138. (New) A method of treating cancer comprising administering to an individual an effective amount of a butyrylcholinesterase variant selected from SEQ ID NOS: 2, 4, 6, 8, 10, 12, 14, 24, 26, 28, 30, 32, 34, 36, 38, 40, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 78, 80, 82, 84, 86, 88, 90, 92, 94, 96, 98, 100, 102, 104, 106, 108, 110, 112, 114, 116, 118, 120, 122, 124, 126, 128, 130, 132, 134, 136, 138, 140, 142, 144,146, 148, 150, 152, 154, 156, 158, 160, 162, 164,166,168, 170, 172, 174, 176, 178, 180, 182, 184, 186, 188, 190, 192, 194, and 196, or functional fragment thereof, exhibiting increased capability to convert a camptothecin derivative to a topoisomerase inhibitor compared to butyrylcholinesterase.

- 139. (New) The method of Claim 138, wherein said cancer is metastatic colorectal cancer.
 - 140. (New) The method of Claim 138, wherein said cancer is ovarian cancer.
 - 141. (New) The method of Claim 138, wherein said cancer is lung cancer.
- 142. (New) The method of Claim 138, wherein said cancer is non-Hodgkin's lymphoma.
 - 143. (New) The method of Claim 138, wherein said topoisomerase inhibitor is SN-38.
- 144. (New) The method of Claim 143, wherein said camptothecin derivative is CPT-11.
- 145. (New) The method of Claim 138, wherein said butyrylcholinesterase variant further comprises an antibody or antibody fragment that specifically binds the epidermal growth factor receptor (EGFR).
- 146. (New) The method of Claim 145, wherein said antibody or antibody fragment comprises an amino acid sequence as shown in SEQ ID NOS: 18 and 20.
- 147. (New) The method of Claim 138, wherein said butyrylcholinesterase variant further comprises an antibody or antibody fragment that specifically binds the CD20 cell surface antigen.
- 148. (New) The method of Claim 147, wherein said antibody or antibody fragment comprises an amino acid sequence as shown in SEQ ID NO: 198.
- 149. (New) The method of Claim 138, wherein said butyrylcholinesterase variant comprises the amino acid sequence designated as SEQ ID NO: 180, or functional fragment thereof.
- 150. (New) The method of Claim 138, wherein said functional fragment is an L530 truncation (SEQ ID NO: 201).